



## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/696,488A  
Source: 1FW0  
Date Processed by STIC: 12/30/04

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PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

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Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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Revised 05/17/04

10/696,488A

SEQUENCE LISTING

<110> Cuenoud, Bernard  
Altmann, Karl-Heinz  
Martin, Pierre  
Moser, Heinz Ernst

These  
are  
prior data  
<120> 2'-Substituted Nucleosides and Oligonucleotide Derivatives  
<130> 4-20890B/C1

<150> <140> 09/753,943  
<151> <141> 2001-01-03

<150> 09/194,844  
<151> 1999-05-14

<150> PCT/EP97/02738  
<151> 1998-05-27

<150> Switzerland 1432/96  
<151> 1996-06-06

delete  
<160> NUMBER OF SEQUENCES: 22  
<170> SOFTWARE: FastSEQ for Windows Version 4.0

<210> SEQ ID NO: 1  
<211> LENGTH: 20  
<212> TYPE: RNA  
<213> ORGANISM: Homo sapiens

<400> SEQ ID NO: 1  
aaugcauguc acaggcggga

20

(sample) submitted file)

Do NOT insert alphabetical  
headings in "New"  
Sequence Ruler  
Sequence Listings. This  
occurs throughout the  
submitted file.